

SEQUENCE LISTING

<110> Beer, Steven V.
Bauer, David W.

<120> OOMYCETE-RESISTANT TRANSGENIC PLANTS BY VIRTUE OF
PATHOGEN-INDUCED EXPRESSION OF A HETEROLOGOUS
HYPERSENSITIVE RESPONSE ELICITOR

<130> 19603/2501

<140>

<141>

<150> 60/178,565

<151> 2000-01-26

<160> 26

<170> PatentIn Ver. 2.1

<210> 1

<211> 338

<212> PRT

<213> *Erwinia chrysanthemi*

<400> 1

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| Met | Gln | Ile | Thr | Ile | Lys | Ala | His | Ile | Gly | Gly | Asp | Leu | Gly | Val | Ser |
| 1 | | | | | 5 | | | | 10 | | | | | 15 | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Leu | Gly | Ala | Gln | Gly | Leu | Lys | Gly | Leu | Asn | Ser | Ala | Ala | Ser | Ser |
| | 20 | | | | | | | 25 | | | | | | 30 | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Gly | Ser | Ser | Val | Asp | Lys | Leu | Ser | Ser | Thr | Ile | Asp | Lys | Leu | Thr |
| | 35 | | | | | | | 40 | | | | | 45 | | |

| | | | | | | | | | | | | | | | |
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| Ser | Ala | Leu | Thr | Ser | Met | Met | Phe | Gly | Gly | Ala | Leu | Ala | Gln | Gly | Leu |
| | 50 | | | | | 55 | | | | | | | 60 | | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Ala | Ser | Ser | Lys | Gly | Leu | Gly | Met | Ser | Asn | Gln | Leu | Gly | Gln | Ser |
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| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Phe | Gly | Asn | Gly | Ala | Gln | Gly | Ala | Ser | Asn | Leu | Leu | Ser | Val | Pro | Lys |
| | | | 85 | | | | | | 90 | | | | | 95 | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Gly | Gly | Asp | Ala | Leu | Ser | Lys | Met | Phe | Asp | Lys | Ala | Leu | Asp | Asp |
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Leu Ala Asn Ser Met Leu Asn Ala Ser Gln Met Thr Gln Gly Asn Met
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Asn Ala Phe Gly Ser Gly Val Asn Asn Ala Leu Ser Ser Ile Leu Gly
145 150 155 160

Asn Gly Leu Gly Gln Ser Met Ser Gly Phe Ser Gln Pro Ser Leu Gly
165 170 175

Ala Gly Gly Leu Gln Gly Leu Ser Gly Ala Gly Ala Phe Asn Gln Leu
180 185 190

Gly Asn Ala Ile Gly Met Gly Val Gly Gln Asn Ala Ala Leu Ser Ala
195 200 205

Leu Ser Asn Val Ser Thr His Val Asp Gly Asn Asn Arg His Phe Val
210 215 220

Asp Lys Glu Asp Arg Gly Met Ala Lys Glu Ile Gly Gln Phe Met Asp
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Gln Tyr Pro Glu Ile Phe Gly Lys Pro Glu Tyr Gln Lys Asp Gly Trp
245 250 255

Ser Ser Pro Lys Thr Asp Asp Lys Ser Trp Ala Lys Ala Leu Ser Lys
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Pro Asp Asp Asp Gly Met Thr Gly Ala Ser Met Asp Lys Phe Arg Gln
275 280 285

Ala Met Gly Met Ile Lys Ser Ala Val Ala Gly Asp Thr Gly Asn Thr
290 295 300

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<211> 2141

<212> DNA

<213> *Erwinia chrysanthemi*

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<210> 3

<211> 403

<212> PRT

<213> *Erwinia amylovora*

<400> 3

09770693-012601

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| Met | Ser | Leu | Asn | Thr | Ser | Gly | Leu | Gly | Ala | Ser | Thr | Met | Gln | Ile | Ser | |
| 1 | | | | | 5 | | | | 10 | | | | | 15 | | |
| Ile | Gly | Gly | Ala | Gly | Gly | Asn | Asn | Gly | Leu | Leu | Gly | Thr | Ser | Arg | Gln | |
| | | | 20 | | | | | 25 | | | | | 30 | | | |
| Asn | Ala | Gly | Leu | Gly | Gly | Asn | Ser | Ala | Leu | Gly | Leu | Gly | Gly | Gly | Asn | |
| | | 35 | | | | | 40 | | | | | 45 | | | | |
| Gln | Asn | Asp | Thr | Val | Asn | Gln | Leu | Ala | Gly | Leu | Leu | Thr | Gly | Met | Met | |
| | 50 | | | | | 55 | | | | | | 60 | | | | |
| Met | Met | Met | Ser | Met | Met | Gly | Gly | Gly | Gly | Leu | Met | Gly | Gly | Gly | Leu | |
| | 65 | | | | 70 | | | | | 75 | | | | | 80 | |
| Gly | Gly | Gly | Leu | Gly | Asn | Gly | Leu | Gly | Gly | Ser | Gly | Gly | Leu | Gly | Glu | |
| | | | 85 | | | | | 90 | | | | | 95 | | | |
| Gly | Leu | Ser | Asn | Ala | Leu | Asn | Asp | Met | Leu | Gly | Gly | Ser | Leu | Asn | Thr | |
| | | | 100 | | | | | 105 | | | | | 110 | | | |
| Leu | Gly | Ser | Lys | Gly | Gly | Asn | Asn | Thr | Thr | Ser | Thr | Thr | Asn | Ser | Pro | |
| | | 115 | | | | 120 | | | | | | | 125 | | | |
| Leu | Asp | Gln | Ala | Leu | Gly | Ile | Asn | Ser | Thr | Ser | Gln | Asn | Asp | Asp | Ser | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Thr | Ser | Gly | Thr | Asp | Ser | Thr | Ser | Asp | Ser | Ser | Asp | Pro | Met | Gln | Gln | |
| 145 | | | | 150 | | | | | 155 | | | | | 160 | | |
| Leu | Leu | Lys | Met | Phe | Ser | Glu | Ile | Met | Gln | Ser | Leu | Phe | Gly | Asp | Gly | |
| | | | 165 | | | | | 170 | | | | | 175 | | | |
| Gln | Asp | Gly | Thr | Gln | Gly | Ser | Ser | Ser | Gly | Gly | Lys | Gln | Pro | Thr | Glu | |
| | | 180 | | | | | | 185 | | | | | 190 | | | |
| Gly | Glu | Gln | Asn | Ala | Tyr | Lys | Lys | Gly | Val | Thr | Asp | Ala | Leu | Ser | Gly | |
| | 195 | | | | | 200 | | | | | 205 | | | | | |
| Leu | Met | Gly | Asn | Gly | Leu | Ser | Gln | Leu | Leu | Gly | Asn | Gly | Gly | Leu | Gly | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Gly | Gly | Gln | Gly | Gly | Asn | Ala | Gly | Thr | Gly | Leu | Asp | Gly | Ser | Ser | Leu | |
| 225 | | | | 230 | | | | | 235 | | | | | 240 | | |
| Gly | Gly | Lys | Gly | Leu | Gln | Asn | Leu | Ser | Gly | Pro | Val | Asp | Tyr | Gln | Gln | |
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Leu Gly Asn Ala Val Gly Thr Gly Ile Gly Met Lys Ala Gly Ile Gln
260 265 270

Ala Leu Asn Asp Ile Gly Thr His Arg His Ser Ser Thr Arg Ser Phe
275 280 285

Val Asn Lys Gly Asp Arg Ala Met Ala Lys Glu Ile Gly Gln Phe Met
290 295 300

Asp Gln Tyr Pro Glu Val Phe Gly Lys Pro Gln Tyr Gln Lys Gly Pro
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Gly Gln Glu Val Lys Thr Asp Asp Lys Ser Trp Ala Lys Ala Leu Ser
325 330 335

Lys Pro Asp Asp Asp Gly Met Thr Pro Ala Ser Met Glu Gln Phe Asn
340 345 350

Lys Ala Lys Gly Met Ile Lys Arg Pro Met Ala Gly Asp Thr Gly Asn
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Gly Asn Leu Gln Ala Arg Gly Ala Gly Gly Ser Ser Leu Gly Ile Asp
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<211> 1288

<212> DNA

<213> *Erwinia amylovora*

<400> 4

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<211> 341

<212> PRT

<213> *Pseudomonas syringae*

<400> 5

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| Ala | Leu | Val | Leu | Val | Arg | Pro | Glu | Ala | Glu | Thr | Thr | Gly | Ser | Thr | Ser |
| | | | 20 | | | | | | 25 | | | | | 30 | |
| Ser | Lys | Ala | Leu | Gln | Glu | Val | Val | Val | Lys | Leu | Ala | Glu | Glu | Leu | Met |
| | | | 35 | | | | | 40 | | | | | | 45 | |
| Arg | Asn | Gly | Gln | Leu | Asp | Asp | Ser | Ser | Pro | Leu | Gly | Lys | Leu | Leu | Ala |
| | | | 50 | | | | | 55 | | | | | | 60 | |
| Lys | Ser | Met | Ala | Ala | Asp | Gly | Lys | Ala | Gly | Gly | Gly | Ile | Glu | Asp | Val |
| | | | 65 | | | 70 | | | | | 75 | | | | 80 |
| Ile | Ala | Ala | Leu | Asp | Lys | Leu | Ile | His | Glu | Lys | Leu | Gly | Asp | Asn | Phe |
| | | | 85 | | | | | | 90 | | | | | 95 | |
| Gly | Ala | Ser | Ala | Asp | Ser | Ala | Ser | Gly | Thr | Gly | Gln | Gln | Asp | Leu | Met |
| | | | 100 | | | | | | 105 | | | | | 110 | |
| Thr | Gln | Val | Leu | Asn | Gly | Leu | Ala | Lys | Ser | Met | Leu | Asp | Asp | Leu | Leu |
| | | | 115 | | | | | 120 | | | | | | 125 | |
| Thr | Lys | Gln | Asp | Gly | Gly | Thr | Ser | Phe | Ser | Glu | Asp | Asp | Met | Pro | Met |
| | | | 130 | | | | | 135 | | | | | 140 | | |
| Leu | Asn | Lys | Ile | Ala | Gln | Phe | Met | Asp | Asp | Asn | Pro | Ala | Gln | Phe | Pro |
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Lys Pro Asp Ser Gly Ser Trp Val Asn Glu Leu Lys Glu Asp Asn Phe
165 170 175

Leu Asp Gly Asp Glu Thr Ala Ala Phe Arg Ser Ala Leu Asp Ile Ile
180 185 190

Gly Gln Gln Leu Gly Asn Gln Gln Ser Asp Ala Gly Ser Leu Ala Gly
195 200 205

Thr Gly Gly Gly Leu Gly Thr Pro Ser Ser Phe Ser Asn Asn Ser Ser
210 215 220

Val Met Gly Asp Pro Leu Ile Asp Ala Asn Thr Gly Pro Gly Asp Ser
225 230 235 240

Gly Asn Thr Arg Gly Glu Ala Gly Gln Leu Ile Gly Glu Leu Ile Asp
245 250 255

Arg Gly Leu Gln Ser Val Leu Ala Gly Gly Gly Leu Gly Thr Pro Val
260 265 270

Asn Thr Pro Gln Thr Gly Thr Ser Ala Asn Gly Gly Gln Ser Ala Gln
275 280 285

Asp Leu Asp Gln Leu Leu Gly Gly Leu Leu Leu Lys Gly Leu Glu Ala
290 295 300

Thr Leu Lys Asp Ala Gly Gln Thr Gly Thr Asp Val Gln Ser Ser Ala
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<210> 6
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<212> DNA
<213> *Pseudomonas syringae*

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<211> 344

<212> PRT

<213> *Pseudomonas solanacearum*

<400> 7

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Val Gln Asp Leu Ile Lys Gln Val Glu Lys Asp Ile Leu Asn Ile Ile
35 40 45

Ala Ala Leu Val Gln Lys Ala Ala Gln Ser Ala Gly Gly Asn Thr Gly
50 55 60

Asn Thr Gly Asn Ala Pro Ala Lys Asp Gly Asn Ala Asn Ala Gly Ala
65 70 75 80

Asn Asp Pro Ser Lys Asn Asp Pro Ser Lys Ser Gln Ala Pro Gln Ser
85 90 95

Ala Asn Lys Thr Gly Asn Val Asp Asp Ala Asn Asn Gln Asp Pro Met
100 105 110

Gln Ala Leu Met Gln Leu Leu Glu Asp Leu Val Lys Leu Leu Lys Ala
115 120 125

Ala Leu His Met Gln Gln Pro Gly Gly Asn Asp Lys Gly Asn Gly Val
130 135 140

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Gly Gly Ala Asn Gly Ala Lys Gly Ala Gly Gly Gln Gly Gly Leu Ala
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Glu Ala Leu Gln Glu Ile Glu Gln Ile Leu Ala Gln Leu Gly Gly Gly
165 170 175

Gly Ala Gly Ala Gly Gly Ala Gly Gly Val Gly Gly Ala Gly Gly
180 185 190

Ala Asp Gly Gly Ser Gly Ala Gly Gly Ala Gly Gly Ala Asn Gly Ala
195 200 205

Asp Gly Gly Asn Gly Val Asn Gly Asn Gln Ala Asn Gly Pro Gln Asn
210 215 220

Ala Gly Asp Val Asn Gly Ala Asn Gly Ala Asp Asp Gly Ser Glu Asp
225 230 235 240

Gln Gly Gly Leu Thr Gly Val Leu Gln Lys Leu Met Lys Ile Leu Asn
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Ala Leu Val Gln Met Met Gln Gln Gly Gly Leu Gly Gly Gly Asn Gln
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Ala Gln Gly Gly Ser Lys Gly Ala Gly Asn Ala Ser Pro Ala Ser Gly
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Ala Asn Pro Gly Ala Asn Gln Pro Gly Ser Ala Asp Asp Gln Ser Ser
290 295 300

Gly Gln Asn Asn Leu Gln Ser Gln Ile Met Asp Val Val Lys Glu Val
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<211> 1035

<212> DNA

<213> *Pseudomonas solanacearum*

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<212> DNA
<213> Solanum tuberosum

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Ser Arg

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 20 25 30

Ser Gln

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